





Sheet 1 of 2

SUPPLEMENTAL INFORMATION DISCLOSURE CITATION	Docket No. 1073.060A	Serial No. 09/832,786
	Applicant: DILLER et al.	
	Filing Date: April 11, 2001	Group: 1631

U.S. PATENT DOCUMENTS								
Examiner Initial		Document Number	Date	Name	Class	Subclass	Filing Date If Appropriate	
CDL	AA	5,854,992	12/29/1998	Shakhnovich et al.	702	27		
FOREIGN PATENT DOCUMENTS								
		DOCUMENT NUMBER	Date	Country	Class	Subclass	Translation	
							Yes	No
CDL	BA	WO 01/97098 A2	20Dec2001	PCT				
Other Documents (including Author, Title, Date, pertinent public. etc.)								
CDL	CA	Welch et al., Hammerhead: Fast, Fully Automated Docking Of Flexible Ligands To Protein Binding Sites, Chemistry & Biology, June 1996, Vol. 3, pp. 449-462						
	CB	Rarey, et al., A Fast Flexible Docking Method Using an Incremental Construction Algorithm, J. Mol. Biol., 1996, 261, pp. 470-489						
	CC	Jones et al., Development and Validation of a Genetic Algorithm for Flexible Docking, J. Mol., Biol., 1997, 267, pp. 727-748						
	CD	Sun et al., CombiDOCK: Structure-based Combinatorial Docking and Library Design, Journal of Computer-Aided Molecular Design, 1998, Vol. 12, pp. 597-604						
	CE	Baxter et al., Flexible Docking Using Tabu Search and an Empirical Estimate of Binding Affinity, PROTEINS: Structure, Function, and Genetics, 1998, 33, pp. 367-382						
	CF	Diller et al., High Throughput Docking for Library Design and Library Prioritization, PROTEINS: Structure, Function, and Genetics, 2001, 43, pp. 113-124						
	CG	Kirkpatrick, et al., Optimization by Simulated Annealing, SCIENCE, Vol. 220, Number 4598, 13 May 1983, pp. 671-680						
	CH	Makino et al., DREAM++: Flexible Docking Program for Virtual Combinatorial Libraries; Journal of Computer-Aided Molecular Design, 13, 513-532, 1999						

Examiner 	Date Considered 10.30.03
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Examiner <i>[Signature]</i>	Date Considered 10. 30. 03
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